

Fri May 16 11:34:30 2003

us-09-551-151a-43.sz1ml8.rapb

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OM protein - protein search, using sw model

Run on: May 16, 2003, 10:37:16 ; Search time 17 seconds
(without alignments)
68.083 Million cell updates/sec

Title: US-09-551-151A-43

Perfect score: 64

Sequence: 1 SPQGIAGQRNFN 12

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 362588 seqs, 96450795 residues

Total number of hits satisfying chosen parameters: 65386

Minimum DB seq length: 0

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Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

Published_Applications_AA:*

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- 2: /cgn2_6/ptodata/1/pubppa/PC1_NEW_PUB.pcp:*
- 3: /cgn2_6/ptodata/1/pubppa/US06_NEW_PUB.pcp:*
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- 12: /cgn2_6/ptodata/1/pubppa/US10_PUBCOMB.pcp:*
- 13: /cgn2_6/ptodata/1/pubppa/US60_NEW_PUB.pcp:*
- 14: /cgn2_6/ptodata/1/pubppa/US60_PUBCOMB.pcp:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match Length	ID	Description
1	42	65.6	9 10 US-09-113-696B-26	Sequence 26, Appl
2	42	65.6	9 10 US-09-816-737-2	Sequence 2, Appl
3	42	65.6	15 9 US-10-133-289-1	Sequence 1, Appl
4	42	65.6	15 10 US-09-113-696B-17	Sequence 17, Appl
5	42	65.6	15 10 US-09-816-737-1	Sequence 1, Appl
6	42	65.6	15 10 US-09-935-417-1	Sequence 1, Appl
7	38	59.4	8 10 US-09-756-283A-28	Sequence 28, Appl
8	37	57.8	8 9 US-09-972-772-3	Sequence 3, Appl
9	37	57.8	8 10 US-09-998-831-25	Sequence 25, Appl
10	37	57.8	8 10 US-09-756-283A-56	Sequence 56, Appl
11	37	57.8	8 12 US-10-001-945-3	Sequence 3, Appl
12	35	54.7	7 10 US-09-113-696B-28	Sequence 28, Appl
13	35	54.7	7 10 US-09-816-737-4	Sequence 4, Appl
14	35	54.7	8 10 US-09-756-283A-30	Sequence 30, Appl
15	32	50.0	7 9 US-09-972-772-4	Sequence 4, Appl
16	32	50.0	7 12 US-10-001-945-4	Sequence 4, Appl
17	30	46.9	6 10 US-09-113-696B-27	Sequence 27, Appl
18	30	46.9	6 10 US-09-816-737-3	Sequence 3, Appl
19	30	46.9	13 9 US-10-132-619-1	Sequence 1, Appl

20	28	43.8	8 10 US-09-756-283A-47	Sequence 47, Appl
21	28	43.8	9 10 US-09-736-076-43	Sequence 43, Appl
22	27	42.2	7 9 US-10-062-710-136	Sequence 136, Appl
23	26	40.6	10 9 US-09-572-404B-4039	Sequence 4039, Appl
24	25	39.1	5 10 US-09-861-708-1	Sequence 1, Appl
25	25	39.1	5 10 US-09-113-696B-30	Sequence 30, Appl
26	25	39.1	5 10 US-09-816-737-6	Sequence 6, Appl
27	25	39.1	5 10 US-09-816-737-8	Sequence 8, Appl
28	25	39.1	5 10 US-09-998-831-27	Sequence 27, Appl
29	25	39.1	8 10 US-09-998-831-26	Sequence 26, Appl
30	25	39.1	8 10 US-09-998-831-28	Sequence 28, Appl
31	25	39.1	8 10 US-09-756-283A-29	Sequence 29, Appl
32	25	39.1	8 10 US-09-756-283A-31	Sequence 31, Appl
33	25	39.1	8 10 US-09-756-283A-57	Sequence 57, Appl
34	25	39.1	10 9 US-09-880-748-5208	Sequence 3208, Appl
35	25	39.1	13 9 US-09-783-931-78	Sequence 78, Appl
36	25	39.1	13 10 US-09-908-322-78	Sequence 78, Appl
37	25	39.1	17 10 US-09-864-761-39966	Sequence 39966, Appl
38	24	37.5	8 10 US-09-756-283A-95	Sequence 95, Appl
39	24	37.5	10 9 US-09-572-404B-4172	Sequence 4172, Appl
40	24	37.5	15 9 US-10-024-918-16	Sequence 16, Appl
41	24	37.5	16 9 US-09-947-925A-36	Sequence 36, Appl
42	24	37.5	17 9 US-09-968-561A-59	Sequence 59, Appl
43	24	37.5	17 10 US-09-947-925A-35	Sequence 35, Appl
44	24	37.5	17 10 US-09-192-854-39	Sequence 39, Appl
45	24	37.5	17 10 US-09-192-854-41	Sequence 41, Appl

ALIGNMENTS

RESULT 1
US-09-113-696B-26
Sequence 26, Application US/09113696B
Patent No. US20020010134A1
GENERAL INFORMATION:
APPLICANT: Bhatnagar, Rajendra S.
APPLICANT: Gough, Craig
TITLE OR INVENTION: PEPTIDE COMPOSITIONS MIMICKING TGF-BETA
FILE REFERENCE: 6510-215C1P2
CURRENT APPLICATION NUMBER: US/09/113, 696B
PRIOR FILING DATE: 1998-07-10
PRIOR APPLICATION NUMBER: 08/742, 256
PRIOR FILING DATE: 1996-10-31
PRIOR APPLICATION NUMBER: 08/431, 954
NUMBER OF SEQ ID NOS: 42
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 26
LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Collagen receptor ligands
US-09-113-696B-26
Query Match
Best Local Similarity 100.0%; Pred. No. 3.3e+05;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Oy 2 POGIAGOR 9
Db 2 POGIAGOR 9
RESULT 2
US-09-816-737-2
Sequence 2, Application US/09816737
Patent No. US20020037853A1
GENERAL INFORMATION:
APPLICANT: Bhatnagar, Rajendra S.

TITLE OF INVENTION: "Synthetic Compounds and Compositions"
TITLE OF INVENTION: With Enhanced Cell Binding"
FILE REFERENCE: 06510223CON2
CURRENT APPLICATION NUMBER: US/09/816,737
CURRENT FILING DATE: 2001-03-23
PRIOR APPLICATION NUMBER: 09/328,347
PRIOR FILING DATE: 1999-06-08
PRIOR APPLICATION NUMBER: 08/859,610
PRIOR FILING DATE: 1997-05-20
PRIOR APPLICATION NUMBER: 08/278,878
PRIOR FILING DATE: 1994-07-22
PRIOR APPLICATION NUMBER: 07/804,782
PRIOR FILING DATE: 1991-12-09
PRIOR APPLICATION NUMBER: 07/393,621
PRIOR FILING DATE: 1989-08-14
NUMBER OF SEQ ID NOS: 14
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: synthetic peptide
US-09-816-737-2

Query Match 65.6%; Score 42; DB 10; Length 9;
Best Local Similarity 100.0%; Pred. No. 3.3e+05;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 POGIAGOR 9
DB 5 POGIAGOR 9

RESULT 3
US-10-133-289-1
Sequence 1, Application US/10133289
Publication No. US20030077825A1
GENERAL INFORMATION:
APPLICANT: Rajendra S. Bhatnagar
APPLICANT: Jing Jing Qian
TITLE OF INVENTION: Structures Useful for Bone Engineering
TITLE OF INVENTION: and Methods
FILE REFERENCE: UCA1224
CURRENT APPLICATION NUMBER: US/10/133,289
CURRENT FILING DATE: 2002-04-25
PRIOR APPLICATION NUMBER: US/09/561,554
PRIOR FILING DATE: 2000-04-28
NUMBER OF SEQ ID NOS: 2
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 15
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Synthetic Peptide
US-10-133-289-1

Query Match 65.6%; Score 42; DB 9; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.2;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 POGIAGOR 9
DB 5 POGIAGOR 12

RESULT 4
US-09-113-696B-17
Sequence 17, Application US/09113696B
Patent No. US20020010134A1
GENERAL INFORMATION:
APPLICANT: Bhatnagar, Rajendra S.

APPLICANT: Qian, Jing Jing
APPLICANT: Gough, Craig
TITLE OF INVENTION: PEPTIDE COMPOSITIONS MIMICKING TGF-BETA
TITLE OF INVENTION: ACTIVITY
FILE REFERENCE: 6510-215CIP2
CURRENT APPLICATION NUMBER: US/09/113,696B
CURRENT FILING DATE: 1998-07-10
PRIOR APPLICATION NUMBER: 08/742,256
PRIOR FILING DATE: 1996-10-31
PRIOR APPLICATION NUMBER: 08/431,954
PRIOR FILING DATE: 1995-05-01
NUMBER OF SEQ ID NOS: 42
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 17
LENGTH: 15
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: TGF-beta mimic
US-09-113-696B-17

Query Match 65.6%; Score 42; DB 10; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.2;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 POGIAGOR 9
DB 5 POGIAGOR 12

RESULT 5
US-09-816-737-1
Sequence 1, Application US/09816737
Patent No. US20020037853A1
GENERAL INFORMATION:
APPLICANT: Bhatnagar, Rajendra S.
TITLE OF INVENTION: "Synthetic Compounds and Compositions"
TITLE OF INVENTION: With Enhanced Cell Binding"
FILE REFERENCE: 06510223CON2
CURRENT APPLICATION NUMBER: US/09/816,737
CURRENT FILING DATE: 2001-03-23
PRIOR APPLICATION NUMBER: 09/328,347
PRIOR FILING DATE: 1999-06-08
PRIOR APPLICATION NUMBER: 08/859,610
PRIOR FILING DATE: 1997-05-20
PRIOR APPLICATION NUMBER: 08/278,878
PRIOR FILING DATE: 1994-07-22
PRIOR APPLICATION NUMBER: 07/804,782
PRIOR FILING DATE: 1991-12-09
PRIOR APPLICATION NUMBER: 07/393,621
PRIOR FILING DATE: 1989-08-14
NUMBER OF SEQ ID NOS: 14
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 15
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: synthetic peptide
US-09-816-737-1

Query Match 65.6%; Score 42; DB 10; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.2;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 POGIAGOR 9
DB 5 POGIAGOR 12

RESULT 6
US-09-935-417-1
Sequence 1, Application US/09935417

Patent No. US20020062145A1
GENERAL INFORMATION:
APPLICANT: Rudakov, Leon V.
APPLICANT: Imran, Mir A.
APPLICANT: Dinh, Linh
APPLICANT: Davidian, Ara
APPLICANT: Laikin, Kevin
TITLE OF INVENTION: Composite Expandable Device with Polymeric Covering and Bioactive
FILE REFERENCE: 52200-8006 US01
CURRENT APPLICATION NUMBER: US/09/935,417
CURRENT FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: US 09/285,691
PRIOR FILING DATE: 1999-08-30
NUMBER OF SEQ ID NOS: 1
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 1
LENGTH: 15
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: portion of a1 chain of collagen
US-09-935-417-1

Query Match 65.6%; Score 42; DB 10; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.2;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 2 POGIAGOR 9
Db 5 POGIAGOR 12

RESULT 7
US-09-756-283A-28
Sequence 28, Application US/09756283A
Patent No. US20020151478A1
GENERAL INFORMATION:
APPLICANT: Chernajovsky, Yuli
APPLICANT: Dreja, Hanna Stina
APPLICANT: Adams, Gillian
TITLE OF INVENTION: Latent Fusion Protein
FILE REFERENCE: 0623.1000000
CURRENT APPLICATION NUMBER: US/09/756,283A
CURRENT FILING DATE: 2001-01-09
NUMBER OF SEQ ID NOS: 100
SOFTWARE: PatentIn version 3.0
SEQ ID NO 28
LENGTH: 8
TYPE: PRT
ORGANISM: Homo sapiens
US-09-756-283A-28

Query Match 59.4%; Score 38; DB 10; Length 8;
Best Local Similarity 87.5%; Pred. No. 3.3e+05;
Matches 7; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

OY 1 SFOGIAGQ 8
Db 1 AFOGIAGQ 8

RESULT 8
US-09-972-772-3
Sequence 3, Application US/09972772
Publication No. US20020193298A1
GENERAL INFORMATION:
APPLICANT: Olson, Gary L.
APPLICANT: Self, Christopher
APPLICANT: Lee, Lily
APPLICANT: Cook, Charles M.
TITLE OF INVENTION: THERAPEUTIC AGENTS AND METHODS OF USE THEREOF FOR THE
TITLE OF INVENTION: MODULATION OF ANGIOGENESIS

FILE REFERENCE: PRT-106CP
CURRENT APPLICATION NUMBER: US/09/972,772
CURRENT FILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: US 09/704,251
PRIOR FILING DATE: 2000-11-01
NUMBER OF SEQ ID NOS: 35
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 3
LENGTH: 8
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Motifs
NAME/KEY: VARIANT
LOCATION: 8
OTHER INFORMATION: Xaa at position 8 represents D-Arginine
US-09-972-772-3

Query Match 57.8%; Score 37; DB 9; Length 8;
Best Local Similarity 100.0%; Pred. No. 3.3e+05;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 2 POGIAGQ 8
Db 1 POGIAGQ 7

RESULT 9
US-09-998-831-25
Sequence 25, Application US/09998831
Patent No. US20020119153A1
GENERAL INFORMATION:
APPLICANT: Philip E. Thorpe
APPLICANT: Rolf A. Breken
TITLE OF INVENTION: ANTIBODY CONJUGATE COMPOSITIONS FOR SELECTIVELY
FILE REFERENCE: 4001.002584
CURRENT APPLICATION NUMBER: US/09/998,831
CURRENT FILING DATE: 2001-11-30
PRIOR APPLICATION NUMBER: 09/561,108
PRIOR FILING DATE: 2000-04-28
NUMBER OF SEQ ID NOS: 44
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 25
LENGTH: 8
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: SYNTHETIC
OTHER INFORMATION: PEPTIDE
US-09-998-831-25

Query Match 57.8%; Score 37; DB 10; Length 8;
Best Local Similarity 100.0%; Pred. No. 3.3e+05;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 2 POGIAGQ 8
Db 2 POGIAGQ 8

RESULT 10
US-09-756-283A-56
Sequence 56, Application US/09756283A
Patent No. US20020151478A1
GENERAL INFORMATION:
APPLICANT: Chernajovsky, Yuli
APPLICANT: Dreja, Hanna Stina
APPLICANT: Adams, Gillian
TITLE OF INVENTION: Latent Fusion Protein
FILE REFERENCE: 0623.1000000
CURRENT APPLICATION NUMBER: US/09/756,283A
CURRENT FILING DATE: 2001-01-09

NUMBER OF SEQ ID NOS: 100
SOFTWARE: PatentIn version 3.0
SEQ ID NO: 56
LENGTH: 8
TYPE: PRT
ORGANISM: Bos taurus
US-09-756-283A-56

Query Match
Best Local Similarity 100.0%; Score 37; DB 10; Length 8;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 POGIAGO 8
Db 2 POGIAGO 8

RESULT 11
US-10-001-945-3
Sequence 3, Application US/10001945
Patent No. US20020151493A1
GENERAL INFORMATION:
APPLICANT: Olson, Gary L.
APPLICANT: Self, Christopher
APPLICANT: Lee, Lily
APPLICANT: Cook, Charles M.
APPLICANT: Birkopf, Jens
TITLE OF INVENTION: THERAPEUTIC AGENTS AND METHODS OF USE THEREOF FOR THE
TITLE OF INVENTION: MODULATION OF ANGIOGENESIS
FILE REFERENCE: PPI-106CP2
CURRENT APPLICATION NUMBER: US/10/001,945
CURRENT FILING DATE: 2001-11-01
PRIOR APPLICATION NUMBER: US 09/972,772
PRIOR FILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: US 09/704,251
PRIOR FILING DATE: 2000-11-01
NUMBER OF SEQ ID NOS: 35
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 3
LENGTH: 8
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Motifs
NAME/KEY: VARIANT
LOCATION: 8
OTHER INFORMATION: Xaa at position 8 represents D-Arginine
US-10-001-945-3

Query Match
Best Local Similarity 100.0%; Score 37; DB 12; Length 8;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 POGIAGO 8
Db 1 POGIAGO 7

RESULT 12
US-09-113-696B-28
Sequence 28, Application US/09113696B
Patent No. US20020010134A1
GENERAL INFORMATION:
APPLICANT: Bhatnagar, Rajendra S.
APPLICANT: Qian, Jing Jiang
APPLICANT: Gough, Craig
TITLE OF INVENTION: PEPTIDE COMPOSITIONS MIMICKING TGF-BETA
FILE REFERENCE: 6510-215CIP2
CURRENT APPLICATION NUMBER: US/09/113,696B
CURRENT FILING DATE: 1998-07-10
PRIOR APPLICATION NUMBER: 08/742,256
PRIOR FILING DATE: 1996-10-31

PRIOR APPLICATION NUMBER: 08/431,954
PRIOR FILING DATE: 1995-05-01
NUMBER OF SEQ ID NOS: 42
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 28
LENGTH: 7
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Collagen receptor ligands
US-09-113-696B-28

Query Match
Best Local Similarity 100.0%; Score 35; DB 10; Length 7;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 OGIAGOR 9
Db 1 OGIAGOR 7

RESULT 13
US-09-816-737-4
Sequence 4, Application US/09816737
Patent No. US20020037853A1
GENERAL INFORMATION:
APPLICANT: Bhatnagar, Rajendra S.
TITLE OF INVENTION: "Synthetic Compounds and Compositions
TITLE OF INVENTION: With Enhanced Cell Binding"
FILE REFERENCE: 06510223CON2
CURRENT APPLICATION NUMBER: US/09/816,737
CURRENT FILING DATE: 2001-03-23
PRIOR APPLICATION NUMBER: 09/328,347
PRIOR FILING DATE: 1999-06-08
PRIOR APPLICATION NUMBER: 08/859,610
PRIOR FILING DATE: 1997-05-20
PRIOR APPLICATION NUMBER: 08/278,878
PRIOR FILING DATE: 1994-07-22
PRIOR APPLICATION NUMBER: 07/804,782
PRIOR FILING DATE: 1991-12-09
PRIOR APPLICATION NUMBER: 07/393,621
PRIOR FILING DATE: 1989-08-14
NUMBER OF SEQ ID NOS: 14
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 4
LENGTH: 7
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: synthetic peptide
US-09-816-737-4

Query Match
Best Local Similarity 100.0%; Score 35; DB 10; Length 7;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 OGIAGOR 9
Db 1 OGIAGOR 7

RESULT 14
US-09-756-283A-30
Sequence 30, Application US/09756283A
Patent No. US20020151478A1
GENERAL INFORMATION:
APPLICANT: Chernajovsky, Yuli
APPLICANT: Dreja, Hanna Stina
APPLICANT: Adams, Gillian
TITLE OF INVENTION: Latent Fusion Protein
FILE REFERENCE: 0623-1000000
CURRENT APPLICATION NUMBER: US/09/756,283A
CURRENT FILING DATE: 2001-01-09

NUMBER OF SEQ ID NOS: 100
SOFTWARE: PatentIn version 3.0
SEQ ID NO 30
LENGTH: 8
TYPE: PRT
ORGANISM: Homo sapiens
US-09-756-283A-30

Query Match 54.7%; Score 35; DB 10; Length 8;
Best Local Similarity 85.7%; Pred. No. 3.3e+05;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 2 POGIAG 8
DB 2 POGIAG 8

RESULT 15
US-09-972-772-4
Sequence 4; Application US/09972772
Publication NO. US20020193298A1
GENERAL INFORMATION:
APPLICANT: Olson, Gary L.
APPLICANT: Self, Christopher
APPLICANT: Lee, Lily
APPLICANT: Cook, Charles M.
TITLE OF INVENTION: THERAPEUTIC AGENTS AND METHODS OF USE THEREOF FOR THE
TITLE OF INVENTION: MODULATION OF ANGIOGENESIS
FILE REFERENCE: PPI-106CP
CURRENT APPLICATION NUMBER: US/09/972,772
CURRENT FILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: US 09/704,251
PRIOR FILING DATE: 2000-11-01
NUMBER OF SEQ ID NOS: 35
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 4
LENGTH: 7
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Motifs
US-09-972-772-4

Query Match 50.0%; Score 32; DB 9; Length 7;
Best Local Similarity 100.0%; Pred. No. 3.3e+05;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 POGIAG 7
DB 1 POGIAG 6

Search completed: May 16, 2003, 10:39:08
Job time : 18 secs

